

SEQUENCE LISTING

<110> Mount Sinai School of Medicine of NYU

<120> TRP8, A TRANSIENT RECEPTOR POTENTIAL CHANNEL EXPRESSED IN TASTE RECEPTOR CELL

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 Inches

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                                   650
               645
Pro Ala Leu Val Tyr Thr Asn Leu Ile Thr Phe Ser Glu Glu Ala Pro
                               665
Leu Arg Thr Gly Leu Glu Asp Leu Gln Asp Leu Asp Ser Leu Asp Thr
                           680
Glu Lys Ser Pro Leu Tyr Gly Leu Gln Ser Arg Val Glu Glu Leu Val
                        695
                                            700
Glu Ala Pro Arg Ala Gln Gly Asp Arg Gly Pro Arg Ala Val Phe Leu
                                        715
                    710
Leu Thr Arg Trp Arg Lys Phe Trp Gly Ala Pro Val Thr Val Phe Leu
                                    730
                725
Gly Asn Val Val Met Tyr Phe Ala Phe Leu Phe Leu Phe Thr Tyr Val
                                745
           740
Leu Leu Val Asp Phe Arg Pro Pro Pro Gln Gly Pro Ser Gly Pro Glu
                           760
Val Thr Leu Tyr Phe Trp Val Phe Thr Leu Val Leu Glu Glu Ile Arg
                       775
                                           780
Gln Gly Phe Phe Thr Asp Glu Asp Thr His Leu Val Lys Lys Phe Thr
                   790
                                        795
Leu Tyr Val Gly Asp Asn Trp Asn Lys Cys Asp Met Val Ala Ile Phe
                                   810
               805
Leu Phe Ile Val Gly Val Thr Cys Arg Met Leu Pro Ser Ala Phe Glu
                               825
           820
Ala Gly Arg Thr Val Leu Ala Met Asp Phe Met Val Phe Thr Leu Arg
                           840
                                                845
Leu Ile His Ile Phe Ala Ile His Lys Gln Leu Gly Pro Lys Ile Ile
                       855
                                            860
Val Val Glu Arg Met Met Lys Asp Val Phe Phe Phe Leu Phe Phe Leu
                   870
                                        875
Ser Val Trp Leu Val Ala Tyr Gly Val Thr Thr Gln Ala Leu Leu His
                885
                                    890
Pro His Asp Gly Arg Leu Glu Trp Ile Phe Arg Arg Val Leu Tyr Arg
                                905
Pro Tyr Leu Gln Ile Phe Gly Gln Ile Pro Leu Asp Glu Ile Asp Glu
                            920
                                                925
Ala Arg Val Asn Cys Ser Thr His Pro Leu Leu Leu Glu Asp Ser Pro
                        935
Ser Cys Pro Ser Leu Tyr Ala Asn Trp Leu Val Ile Leu Leu Val
                    950
                                        955
Thr Phe Leu Leu Val Thr Asn Val Leu Leu Met Asn Leu Leu Ile Ala
                                    970
Met Phe Ser Tyr Thr Phe Gln Val Gln Gly Asn Ala Asp Met Phe
                                985
           980
Trp Lys Phe Gln Arg Tyr Asn Leu Ile Val Glu Tyr His Glu Arg Pro
                           1000
Ala Leu Ala Pro Pro Phe Ile Leu Leu Ser His Leu Ser Leu Thr Leu
                                            1020
                       1015
Arg Arg Val Phe Lys Lys Glu Ala Glu His Lys Arg Glu His Leu Glu
1025
                   1030
                                        1035
Arg Asp Leu Pro Asp Pro Leu Asp Gln Lys Val Val Thr Trp Glu Thr
                1045
                                    1050
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Val	Gln	Lys	Glu	Asn	Phe	Leu	Ser			Glu	Lys	Arg	Arg		Asp
	1060					1065					1070				
Ser	Glu	Gly	Glu	Val	Leu	Arg	Lys	Thr	Ala	His	Arg	Val	Asp	Phe	Ile
1075						1080					1085				
Ala	Lys	Tyr	Leu	Gly	Gly	Leu	Arg	Glu	Gln	Glu	Lys	Arg	Ile	Lys	Cys
1090						1095					1100)			
Leu	Glu	Ser	Gln	Ile	Asn	Tyr	Cys	Ser	Val	Leu	Val	Ser	Ser	Val	Ala
1105					1110	10			1115						1120
Asp	Val	Leu	Ala	Gln	Gly	Gly	Gly	Pro	Arg	Ser	Ser	Gln	His	Cys	Gly
1125					1130					1135					
C1															
GIU	Gly	Ser	Gln	Leu	Val	Ala	Ala	Asp	His	Arg	Gly	Gly	Leu	Asp	Gly
GIU	Gly	Ser	Gln 1140		Val	Ala	Ala	Asp 1145		Arg	Gly	Gly	Leu 1150		Gly
			1140)		Ala Gly		1145	5						Gly

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